

GenCore version 4.5
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OM protein - protein search, using sw model

Run on: July 9, 2002, 12:18:25 ; Search time 106.06 Seconds
(without alignments)
16.593 Million cell updates/sec

Title: US-09-759-484-3
Perfect score: 22
Sequence: 1 AMVSE 5

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 3502263 seqs, 351980561 residues

Total number of hits satisfying chosen parameters: 3502263

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%

Listing first 45 summaries

Database :

1: /cgn2_6/ptodata/2/paa/PCTUS.COMB.pep.*
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3: /cgn2_6/ptodata/2/paa/US07.COMB.pep.*
4: /cgn2_6/ptodata/2/paa/US08.COMB.pep.*
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19: /cgn2_6/ptodata/2/paa/US095.COMB.pep.*
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26: /cgn2_6/ptodata/2/paa/US60.COMB.pep.*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the change being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	22	100.0	5	21	US-09-759-484-3
2	22	100.0	11	21	US-09-759-484-3
3	22	100.0	25	21	US-09-759-484-6
4	22	100.0	40	18	US-09-480-993-14
5	22	100.0	40	24	US-10-044-967-14
6	22	100.0	78	18	US-09-417-507-40089
7	22	100.0	88	22	US-09-834-366-16299

8	22	100.0	88	26	US-60-197-873-16299	Sequence 16299, A
9	22 <th>100.0</th> <td>97</td> <td>20</td> <td>US-09-620-394B-4378</td> <td>Sequence 4378, Ap</td>	100.0	97	20	US-09-620-394B-4378	Sequence 4378, Ap
10	22 <th>100.0</th> <td>98</td> <td>19</td> <td>US-09-595-298A-306</td> <td>Sequence 306, App</td>	100.0	98	19	US-09-595-298A-306	Sequence 306, App
11	22 <th>100.0</th> <td>109</td> <td>19</td> <td>US-09-540-236-2546</td> <td>Sequence 2546, Ap</td>	100.0	109	19	US-09-540-236-2546	Sequence 2546, Ap
12	22 <th>100.0</th> <td>120</td> <td>20</td> <td>US-09-673-840A-163</td> <td>Sequence 163, App</td>	100.0	120	20	US-09-673-840A-163	Sequence 163, App
13	22 <th>100.0</th> <td>121</td> <td>20</td> <td>US-09-675-784A-9227</td> <td>Sequence 9227, Ap</td>	100.0	121	20	US-09-675-784A-9227	Sequence 9227, Ap
14	22 <th>100.0</th> <td>149</td> <td>1</td> <td>PCT-US01-08656-6479</td> <td>Sequence 6475, Ap</td>	100.0	149	1	PCT-US01-08656-6479	Sequence 6475, Ap
15	22 <th>100.0</th> <td>149</td> <td>1</td> <td>PCT-US01-14827-10033</td> <td>Sequence 10033, A</td>	100.0	149	1	PCT-US01-14827-10033	Sequence 10033, A
16	22 <th>100.0</th> <td>149</td> <td>21</td> <td>US-09-760-484-519</td> <td>Sequence 519, App</td>	100.0	149	21	US-09-760-484-519	Sequence 519, App
17	22 <th>100.0</th> <td>152</td> <td>20</td> <td>US-09-620-394B-4377</td> <td>Sequence 4377, Ap</td>	100.0	152	20	US-09-620-394B-4377	Sequence 4377, Ap
18	22 <th>100.0</th> <td>153</td> <td>19</td> <td>US-09-595-298A-305</td> <td>Sequence 305, App</td>	100.0	153	19	US-09-595-298A-305	Sequence 305, App
19	22 <th>100.0</th> <td>155</td> <td>20</td> <td>US-09-620-394B-4376</td> <td>Sequence 304, App</td>	100.0	155	20	US-09-620-394B-4376	Sequence 304, App
20	22 <th>100.0</th> <td>156</td> <td>19</td> <td>US-09-595-298A-304</td> <td>Sequence 39318, A</td>	100.0	156	19	US-09-595-298A-304	Sequence 39318, A
21	22 <th>100.0</th> <td>167</td> <td>20</td> <td>US-09-614-150-39318</td> <td>Sequence 29656, A</td>	100.0	167	20	US-09-614-150-39318	Sequence 29656, A
22	22 <th>100.0</th> <td>167</td> <td>26</td> <td>US-60-173-464-29696</td> <td>Sequence 38944, A</td>	100.0	167	26	US-60-173-464-29696	Sequence 38944, A
23	22 <th>100.0</th> <td>167</td> <td>26</td> <td>US-60-191-637-38944</td> <td>Sequence 30178, A</td>	100.0	167	26	US-60-191-637-38944	Sequence 30178, A
24	22 <th>100.0</th> <td>167</td> <td>26</td> <td>US-60-191-681-30178</td> <td>Sequence 30178, A</td>	100.0	167	26	US-60-191-681-30178	Sequence 30178, A
25	22 <th>100.0</th> <td>186</td> <td>1</td> <td>PCT-US01-18569-2936</td> <td>Sequence 2936, Ap</td>	100.0	186	1	PCT-US01-18569-2936	Sequence 2936, Ap
26	22 <th>100.0</th> <td>192</td> <td>21</td> <td>US-09-721-756-4</td> <td>Sequence 4, Appl1</td>	100.0	192	21	US-09-721-756-4	Sequence 4, Appl1
27	22 <th>100.0</th> <td>210</td> <td>16</td> <td>US-09-248-796-15009</td> <td>Sequence 15009, A</td>	100.0	210	16	US-09-248-796-15009	Sequence 15009, A
28	22 <th>100.0</th> <td>210</td> <td>26</td> <td>US-60-096-409-15009</td> <td>Sequence 524, App</td>	100.0	210	26	US-60-096-409-15009	Sequence 524, App
29	22 <th>100.0</th> <td>212</td> <td>21</td> <td>US-09-760-484-524</td> <td>Sequence 11628, A</td>	100.0	212	21	US-09-760-484-524	Sequence 11628, A
30	22 <th>100.0</th> <td>238</td> <td>24</td> <td>US-10-015-127-11628</td> <td>Sequence 8907, Ap</td>	100.0	238	24	US-10-015-127-11628	Sequence 8907, Ap
31	22 <th>100.0</th> <td>250</td> <td>18</td> <td>US-09-489-039A-8907</td> <td>Sequence 11126, A</td>	100.0	250	18	US-09-489-039A-8907	Sequence 11126, A
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33	22 <th>100.0</th> <td>255</td> <td>16</td> <td>US-60-324-109-24355</td> <td>Sequence 368, App</td>	100.0	255	16	US-60-324-109-24355	Sequence 368, App
34	22 <th>100.0</th> <td>269</td> <td>26</td> <td>US-09-760-484-524</td> <td>Sequence 21673, A</td>	100.0	269	26	US-09-760-484-524	Sequence 21673, A
35	22 <th>100.0</th> <td>282</td> <td>21</td> <td>US-09-252-691C-11126</td> <td>Sequence 7586, Ap</td>	100.0	282	21	US-09-252-691C-11126	Sequence 7586, Ap
36	22 <th>100.0</th> <td>293</td> <td>16</td> <td>US-09-252-691C-11126</td> <td>Sequence 12733, A</td>	100.0	293	16	US-09-252-691C-11126	Sequence 12733, A
37	22 <th>100.0</th> <td>299</td> <td>16</td> <td>US-09-252-691C-11126</td> <td>Sequence 4744, Ap</td>	100.0	299	16	US-09-252-691C-11126	Sequence 4744, Ap
38	22 <th>100.0</th> <td>299</td> <td>16</td> <td>US-09-252-691C-11126</td> <td>Sequence 641, App</td>	100.0	299	16	US-09-252-691C-11126	Sequence 641, App
39	22 <th>100.0</th> <td>299</td> <td>16</td> <td>US-09-252-691C-11126</td> <td>Sequence 21932, A</td>	100.0	299	16	US-09-252-691C-11126	Sequence 21932, A
40	22 <th>100.0</th> <td>314</td> <td>22</td> <td>US-09-897-516-4744</td> <td>Sequence 6, Appl1</td>	100.0	314	22	US-09-897-516-4744	Sequence 6, Appl1
41	22 <th>100.0</th> <td>314</td> <td>22</td> <td>US-09-897-516-4744</td> <td>Sequence 1, Appl1</td>	100.0	314	22	US-09-897-516-4744	Sequence 1, Appl1
42	22 <th>100.0</th> <td>320</td> <td>22</td> <td>US-09-855-708-641</td> <td></td>	100.0	320	22	US-09-855-708-641	
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45	22 <th>100.0</th> <td>346</td> <td>21</td> <td>US-09-787-923-1</td> <td></td>	100.0	346	21	US-09-787-923-1	

ALIGNMENTS

RESULT 1
US-09-759-484-3 Application US/09759484
Sequence 3, Appl1
GENERAL INFORMATION:
APPLICANT: William Harvey Research Limited
TITLE OR INVENTION: Anti-inflammatory compounds
FILE REFERENCE: P019602MO
CURRENT APPLICATION NUMBER: US/09/759,484
CURRENT FILING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 8
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 3
LENGTH: 5
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: synthetic
US-09-759-484-3

Query Match 100.0%; Score 22; DB 21; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.2e+06;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMVSE 5
DB 1 AMVSE 5

RESULT 2
US-09-759-484-4
; Sequence 4, Application US/09759484
; GENERAL INFORMATION:
; APPLICANT: William Harvey Research Limited
; TITLE OF INVENTION: Anti-Inflammatory compounds
; FILE REFERENCE: P019602WO
; CURRENT APPLICATION NUMBER: US/09/759,484
; CURRENT FILING DATE: 2001-01-12
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 4
; LENGTH: 11
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: synthetic
US-09-759-484-4

Query Match 100.0%; Score 22; DB 21; Length 11;
Best Local Similarity 100.0%; Pred. No. 23;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMVSE 5
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Db 1 AMVSE 5

RESULT 3
US-09-759-484-6
; Sequence 6, Application US/09759484
; GENERAL INFORMATION:
; APPLICANT: William Harvey Research Limited
; TITLE OF INVENTION: Anti-Inflammatory compounds
; FILE REFERENCE: P019602WO
; CURRENT APPLICATION NUMBER: US/09/759,484
; CURRENT FILING DATE: 2001-01-12
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 6
; LENGTH: 25
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: synthetic
US-09-759-484-6

Query Match 100.0%; Score 22; DB 21; Length 25;
Best Local Similarity 100.0%; Pred. No. 61;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMVSE 5
| | | | |
Db 1 AMVSE 5

RESULT 4
US-09-480-993-14
; Sequence 14, Application US/09480993
; GENERAL INFORMATION:
; APPLICANT: Shokat, Keyan M.
; TITLE OF INVENTION: High Affinity Kinase Inhibitors for Target Validation
; FILE REFERENCE: 51538-5001-US
; CURRENT APPLICATION NUMBER: US/09/480,993
; CURRENT FILING DATE: 2000-01-11
; EARLIER APPLICATION NUMBER: US 60/115,340
; EARLIER FILING DATE: 1999-01-11
; NUMBER OF SEQ ID NOS: 20

; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 14
; LENGTH: 40
; TYPE: PRT
; ORGANISM: Rous sarcoma virus
; FEATURE:
; OTHER INFORMATION: v-Src kinase
US-09-480-993-14

Query Match 100.0%; Score 22; DB 18; Length 40;
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMVSE 5
| | | | |
Db 10 AMVSE 14

RESULT 5
US-10-044-967-14
; Sequence 14, Application US/10044967
; GENERAL INFORMATION:
; APPLICANT: Shokat, Keyan M.
; TITLE OF INVENTION: High Affinity Kinase Inhibitors for Target Validation
; FILE REFERENCE: 51538-5001-US
; CURRENT APPLICATION NUMBER: US/10/044,967
; CURRENT FILING DATE: 2002-01-15
; PRIOR APPLICATION NUMBER: 09/480,993
; PRIOR FILING DATE: 2000-01-11
; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/115,340
; PRIOR FILING DATE: EARLIER FILING DATE: 1999-01-11
; NUMBER OF SEQ ID NOS: 20
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 14
; LENGTH: 40
; TYPE: PRT
; ORGANISM: Rous sarcoma virus
; FEATURE:
; OTHER INFORMATION: v-Src kinase
US-10-044-967-14

Query Match 100.0%; Score 22; DB 24; Length 40;
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMVSE 5
| | | | |
Db 10 AMVSE 14

RESULT 6
US-09-417-507-40089
; Sequence 40089, Application US/09417507
; GENERAL INFORMATION:
; APPLICANT: KEITH G. WEINSTOCK ET AL.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ASPERGILLUS
; FILE REFERENCE: PAT99-10
; CURRENT APPLICATION NUMBER: US/09/417,507
; CURRENT FILING DATE: 1999-10-14
; NUMBER OF SEQ ID NOS: 44312
; SEQ ID NO 40089
; LENGTH: 78
; TYPE: PRT
; ORGANISM: A.fumigatus
US-09-417-507-40089

Query Match 100.0%; Score 22; DB 18; Length 78;
Best Local Similarity 100.0%; Pred. No. 2.3e+02;

Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMVSE 5
|||||

Db 59 AMVSE 63

RESULT 7

US-09-834-366-16299
; Sequence 16299, Application US/09834366
; GENERAL INFORMATION:
; APPLICANT: Benjamin, Stephane
; APPLICANT: Tanaka, Hiroaki
; APPLICANT: Dumas Milne Edwards, Jean Baptiste
; APPLICANT: Jobert, Severin
; APPLICANT: Giordano, Jean-Yves
; TITLE OF INVENTION: ESTs and Encoded Human Proteins.
; FILE REFERENCE: 81.US2.REG
; CURRENT APPLICATION NUMBER: US/09/834,366
; CURRENT FILING DATE: 2001-04-13
; PRIOR APPLICATION NUMBER: US 60/197,873
; PRIOR FILING DATE: 2000-04-18
; NUMBER OF SEQ ID NOS: 52153
; SOFTWARE: Patent.pm
; SEQ ID NO 16299
; LENGTH: 88
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-834-366-16299

Query Match

Best Local Similarity 100.0%; Score 22; DB 22; Length 88;
Pred. No. 2.7e+02;

Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMVSE 5
|||||

Db 2 AMVSE 6

RESULT 8

US-60-197-873-16299
; Sequence 16299, Application US/60197873
; GENERAL INFORMATION:
; APPLICANT: Benjamin, Stephane
; APPLICANT: Tanaka, Hiroaki
; APPLICANT: Dumas Milne Edwards, Jean Baptiste
; APPLICANT: Jobert, Severin
; APPLICANT: Giordano, Jean-Yves
; TITLE OF INVENTION: ESTs and Encoded Human Proteins.
; FILE REFERENCE: 81.US1.PRO
; CURRENT APPLICATION NUMBER: US/60/197,873
; CURRENT FILING DATE: 2000-04-18
; NUMBER OF SEQ ID NOS: 52153
; SOFTWARE: Patent.pm
; SEQ ID NO 16299
; LENGTH: 88
; TYPE: PRT
; ORGANISM: Homo sapiens
US-60-197-873-16299

Query Match

Best Local Similarity 100.0%; Score 22; DB 26; Length 88;
Pred. No. 2.7e+02;

Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMVSE 5
|||||

Db 2 AMVSE 6

RESULT 9

US-09-620-394B-4378

; Sequence 4378, Application US/09620394B

; GENERAL INFORMATION:
; APPLICANT: ALEXANDROV, Nikolai
; APPLICANT: BROVER, Vyacheslav
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptide
; FILE REFERENCE: 2750-1067P
; CURRENT APPLICATION NUMBER: US/09/620,394B
; CURRENT FILING DATE: 2000-07-21
; NUMBER OF SEQ ID NOS: 9131
; SEQ ID NO 4378
; LENGTH: 97
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: 1..97
; OTHER INFORMATION: Xaa is any amino acid
; NAME/KEY: misc_feature
; LOCATION: 1..97
; OTHER INFORMATION: Ceres Seq. ID 1389404
US-09-620-394B-4378

Query Match

Best Local Similarity 100.0%; Score 22; DB 20; Length 97;
Pred. No. 3e+02;

Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMVSE 5
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Db 75 AMVSE 79

RESULT 10

US-09-595-298A-306
; Sequence 306, Application US/09595298A
; GENERAL INFORMATION:
; APPLICANT: ALEXANDROV, Nikolai
; APPLICANT: BROVER, Vyacheslav
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptide
; FILE REFERENCE: 2750-0953P
; CURRENT APPLICATION NUMBER: US/09/595,298A
; CURRENT FILING DATE: 2000-06-16
; NUMBER OF SEQ ID NOS: 2756
; SOFTWARE: Patent version 3.0
; SEQ ID NO 306
; LENGTH: 98
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
; FEATURE:
; NAME/KEY: peptide
; LOCATION: (1)..(98)
; OTHER INFORMATION: Ceres Seq. ID no. 1023991
; NAME/KEY: misc_feature
; LOCATION: (1)..(1)
; OTHER INFORMATION: Xaa is any aa, unknown or other
US-09-595-298A-306

Query Match

Best Local Similarity 100.0%; Score 22; DB 19; Length 98;
Pred. No. 3.1e+02;

Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMVSE 5
|||||

Db 75 AMVSE 79

RESULT 11

US-09-540-236-2546
; Sequence 2546, Application US/09540236
; GENERAL INFORMATION:

APPLICANT: Gary L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO MORAXELLA CATAR
FILE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2005-001
CURRENT APPLICATION NUMBER: US/09/540,236
CURRENT FILING DATE: 2000-04-04
NUMBER OF SEQ ID NOS: 3840
SEQ ID NO 2546
LENGTH: 109
TYPE: PRT
ORGANISM: M.catarrhalis
US-09-340-236-2546

Query Match 100.0%; Score 22; DB 19; Length 109;
Best Local Similarity 100.0%; Pred. No. 3.5e+02;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 AMVSE 5
Db 25 AMVSE 29

RESULT 12
US-09-673-840A-163
Sequence 163, Application US/09673840A
GENERAL INFORMATION:
APPLICANT: SPECHT, THOMAS
APPLICANT: HINZMANN, BERND
APPLICANT: SCHMITT, ARMIN
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: DAHL, EDGAR
APPLICANT: ROSENTHAL, ANDRE
TITLE OF INVENTION: HUMAN NUCLEIC ACID SEQUENCES FROM NORMAL BLADDER TISSUE
FILE REFERENCE: ALBRE 5
CURRENT APPLICATION NUMBER: US/09/673,840A
CURRENT FILING DATE: 2000-10-23
NUMBER OF SEQ ID NOS: 433
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 163
LENGTH: 120
TYPE: PRT
ORGANISM: Homo sapiens
US-09-673-840A-163

Query Match 100.0%; Score 22; DB 20; Length 120;
Best Local Similarity 100.0%; Pred. No. 3.9e+02;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 AMVSE 5
Db 29 AMVSE 33

RESULT 13
US-09-675-784A-9227
Sequence 9227, Application US/09675784A
GENERAL INFORMATION:
APPLICANT: HARE, ROBERTA S.
APPLICANT: SHAW, KAREN J.
APPLICANT: KESSLER, MARCO
APPLICANT: NOLLING, JORK
APPLICANT: ZENG, QIANDONG
APPLICANT: GREENE, JONATHAN R.
TITLE OF INVENTION: ASPERGILLUS FUNIGATUS NUCLEIC ACIDS AND POLYPEPTIDES,
FILE REFERENCE: 2976-4020051
CURRENT APPLICATION NUMBER: US/09/675,784A
CURRENT FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: 60/156,338
PRIOR FILING DATE: 1999-09-29

NUMBER OF SEQ ID NOS: 13925
SEQ ID NO 9227
LENGTH: 121
TYPE: PRT
ORGANISM: Aspergillus fumigatus
US-09-675-784A-9227

Query Match 100.0%; Score 22; DB 20; Length 121;
Best Local Similarity 100.0%; Pred. No. 3.9e+02;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 AMVSE 5
Db 102 AMVSE 106

RESULT 14
PCT-US01-08656-6479
Sequence 6479, Application PC/TUS0108656
GENERAL INFORMATION:
APPLICANT: Hyseq, Inc
FILE REFERENCE: 21272-066
CURRENT APPLICATION NUMBER: PCT/US01/08656
CURRENT FILING DATE: 2001-04-16
PRIOR APPLICATION NUMBER: 09/522,929
PRIOR FILING DATE: 2000-04-16
PRIOR APPLICATION NUMBER: 09/770,160
PRIOR FILING DATE: 2001-01-26
NUMBER OF SEQ ID NOS: 10994
SOFTWARE: Custom
SEQ ID NO 6479
LENGTH: 149
TYPE: PRT
ORGANISM: Homo sapiens
PCT-US01-08656-6479

Query Match 100.0%; Score 22; DB 1; Length 149;
Best Local Similarity 100.0%; Pred. No. 5e+02;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 AMVSE 5
Db 60 AMVSE 64

RESULT 15
PCT-US01-14827-10033
Sequence 10033, Application PC/TUS0114827
GENERAL INFORMATION:
APPLICANT: Hyseq, Inc
TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
FILE REFERENCE: 21272-104
CURRENT APPLICATION NUMBER: PCT/US01/14827
CURRENT FILING DATE: 2001-05-16
PRIOR APPLICATION NUMBER: 09/577,408
PRIOR FILING DATE: 2000-05-18
NUMBER OF SEQ ID NOS: 16102
SOFTWARE: Custom
SEQ ID NO 10033
LENGTH: 149
TYPE: PRT
ORGANISM: Homo sapiens
PCT-US01-14827-10033

Query Match 100.0%; Score 22; DB 1; Length 149;
Best Local Similarity 100.0%; Pred. No. 5e+02;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 AMVSE 5

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us-09-759-484-3.rapm

Page 5

|||||
Db 60 AMVSE 64

Search completed: July 9, 2002, 12:22:07
Job time: 222 sec

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11. The following table shows the results of the survey of the
12. The following table shows the results of the survey of the
13. The following table shows the results of the survey of the